



Tue Jan 8 08:23:29 2002

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Page 2

[illegible]

Query Match	100.0%	Score 1277	Db 5	Length 618
Best Local Similarity	100.0%	Pred. no. 1, 2e-308		
Matches 618	Conservative	0	Mismatches 0	Indels 0
			Gaps 0	
<p> <b>TELEFAX:</b> (415) 936-7249  <b>INITIALS:</b> J  <b>SEQUENCE CHARACTERISTICS:</b> 2:  <b>LENGTH:</b> 618 amino acids  <b>STRANDEDNESS:</b> single  <b>STRANDEDNESS:</b> stable  <b>TOPOLOGY:</b> linear  <b>MOLECULE TYPE:</b> Protein  <b>PROTID:</b> P08002         </p>				
01	1 HNTAQSGLDQPSYQKAKIEEDSTLSDMNINQKMDICDLSIAHSTSTPQAYG			
02	1 IHNKQGLDQPSYQKAKIEEDSTLSDMNINQKMDICDLSIAHSTSTPQAYG			
03	1 HNTAQSGLDQPSYQKAKIEEDSTLSDMNINQKMDICDLSIAHSTSTPQAYG			
04	6 PVSERISALQPTVYQGNKLEEDSTLSDMNINQKMDICDLSIAHSTSTPQAYG			
05	6 PVSERISALQPTVYQGNKLEEDSTLSDMNINQKMDICDLSIAHSTSTPQAYG			
06	6 PVSERISALQPTVYQGNKLEEDSTLSDMNINQKMDICDLSIAHSTSTPQAYG			
07	121 LQTSKSTSNHNSKALSPPTLEKSLSCSTSLSPNLSKSNALSSSTNPSTLA			
08	121 LQTSKSTSNHNSKALSPPTLEKSLSCSTSLSPNLSKSNALSSSTNPSTLA			
09	121 LQTSKSTSNHNSKALSPPTLEKSLSCSTSLSPNLSKSNALSSSTNPSTLA			
10	121 LQTSKSTSNHNSKALSPPTLEKSLSCSTSLSPNLSKSNALSSSTNPSTLA			
11	121 LQTSKSTSNHNSKALSPPTLEKSLSCSTSLSPNLSKSNALSSSTNPSTLA			
12	121 LQTSKSTSNHNSKALSPPTLEKSLSCSTSLSPNLSKSNALSSSTNPSTLA			
13	181 MSTREKATLTTHMHTLTSPSLAAGCTYTGQGGHACAGCTSLWPPQKANSR			
14	181 MSTREKATLTTHMHTLTSPSLAAGCTYTGQGGHACAGCTSLWPPQKANSR			
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16	181 MSTREKATLTTHMHTLTSPSLAAGCTYTGQGGHACAGCTSLWPPQKANSR			
17	241 HIRNPQKPTLHNSLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTST			
18	241 HIRNPQKPTLHNSLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTST			
19	241 HIRNPQKPTLHNSLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTST			
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22	300 NDQVQKPCDQKQKNSGSDGAWHNSPKPSEFLIMKQSPQVQDQPHLLEQL			
23	300 NDQVQKPCDQKQKNSGSDGAWHNSPKPSEFLIMKQSPQVQDQPHLLEQL			
24	300 NDQVQKPCDQKQKNSGSDGAWHNSPKPSEFLIMKQSPQVQDQPHLLEQL			
25	360 STGVDVQGGDPIHJHGGSSSGLQVAPVYVSSALNDQVQSGKLTAT			
26	360 STGVDVQGGDPIHJHGGSSSGLQVAPVYVSSALNDQVQSGKLTAT			
27	420 GERTVYQVYSLALMDERKEERKQALNKAQSSQSLIRHMAHQLCPALP			
28	420 GERTVYQVYSLALMDERKEERKQALNKAQSSQSLIRHMAHQLCPALP			
29	420 GERTVYQVYSLALMDERKEERKQALNKAQSSQSLIRHMAHQLCPALP			
30	480 GERTVYQVYSLALMDERKEERKQALNKAQSSQSLIRHMAHQLCPALP			
31	480 GERTVYQVYSLALMDERKEERKQALNKAQSSQSLIRHMAHQLCPALP			
32	540 HIRNPQKPTLHNSLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTST			
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54	540 HIRNPQKPTLHNSLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTSTLTST			

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STREET : 225 Franklin Street  
CITY : Boston  
STATE : MA  
COUNTRY : USA  
COMPUTER RELEVABLE FORM :  
MEDIUM TYPE : Floppy disk  
OPERATING SYSTEM : PC-DOS/MS-DOS  
SOFTWARE : Research Release #1.0, Version #1.30  
APPLICATION NUMBER : US/08/511.485  
FILING DATE : 04-JAN-1995  
ATTORNEY/AGENT INFORMATION :  
NAME : CLARK, PAUL T.  
ADDRESS : 100 N. STATE STREET, BOX 30, 462  
BOSTON, MA 02111-0462  
TELEPHONE : 617/542-8900  
TELECOMMUNICATION INFORMATION :  
REFERENCE/BOOKER NUMBER : 075-0/002001  
TELEFAX : 617/542-8906  
FAX : 605154  
INFORMATION FURNISHED TO NO. 8 :  
SEQUENCE CHARACTERISTICS :  
LENGTH : 618 amino acids  
STRANDEDNESS : not relevant  
COMMENTS : both N-terminus and C-terminus  
are of the type : proline

[illegible]

Db	Qy	Db
541	KCPICNGIKGVRTFELS 618	601
VADNNKVIPTFEDSLSLEQRLRIQSEKRCVCKKEDSVFVPCGHVYQCEAFSLR 600	////////////////////	601
	KCPICNGIKGVRTFELS 618	601
	////////////////////	601
	KCPICNGIKGVRTFELS 618	601

05-09-21-971-8  
Sequence B, Application US/00212971B  
GENERAL INFORMATION  
APPLICANT: Korbelik, Robert G  
APPLICANT: Korbelik, Robert G  
APPLICANT: Litton, Reed  
APPLICANT: Litton, Reed  
APPLICANT: Pratt, Christine  
TITLE OF INVENTION: DETECTION AND MODULATION OF TARE AND  
TITLE OF INVENTION: THE DIAMONDS AND THERMOSTAT OF PROLIFERATIVE  
FILE REFERENCE: 07891/069002  
CURRENT PILING DATE: 1998-12-16  
BAILLER APPLICATION NUMBER: 66/017 344  
BAILLER PILING DATE: 1996-11-14  
BAILLER APPLICATION NUMBER: 66/030 580  
BAILLER PILING DATE: 1997-02-13  
NUMBER OF SEQ IN MOS: 17  
SEQ ID NO: 8  
FILED FOR WINDOWS VERSION 4.0  
LENGTH: 618  
OCCASION: ROMO softpicks  
05-09-21-971-8

[illegible]







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RESULTS: 10  
 CSD00000294.4  
 Sequence ID: Application 075/08809256  
 Patent No. 633437  
 GENERAL INFORMATION  
 APPLICANT: Lister, Peter  
 APPLICANT: Macdonald, Alexander E  
 APPLICANT: Macdonald, Peter  
 APPLICANT: Tzeng, Benjamin X  
 APPLICANT: Pratt, Christine  
 TITLE OF INVENTION: 19S AND 49S FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE  
 TITLE OF INVENTION: DISMSE  
 NUMBER OF SEQUENCES: 17  
 NUMBER OF SEQUENCES: 17  
 ADDRESSER: Clark & Ebling LLP  
 STREET: 106 Federal Street  
 STATE: MA  
 COUNTRY: USA  
 ZIP: 02130  
 COMPUTER PROGRAMS FROM:  
 COMPUTER: IBM Compatible  
 MEDIUM TYPE: Database  
 SOFTWARE: PARASOFT FOR WINDOWS VERSION 2.0  
 CURRENT APPLICATION DATA:  
 FILING DATE: 13 FEB 1997  
 CLASSIFICATION: 424  
 PRIOR APPLICATION DATA:  
 FILING DATE: 66/070,590  
 APPLICATION NUMBER: 66/071,254  
 APPLICATION NUMBER: 66/071,254  
 FILING DATE: 14 NOV 1995  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Baker-Brody, Kristina  
 RESIDENCE/COUNTRY NUMBER: 07891/09901  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: 617-425-0200  
 TELEPHONE: 617-425-0405  
 FAX:  
 INFORMATION FOR SEQ ID NO: 14:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 612 amino acids  
 TYPE: amino acid  
 STRANDNESS: single  
 MOLECULE TYPE: protein

[illegible][illegible]

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SRD ID NO 14  
LEADIN: 612  
TITLE: PART TWO  
SUBTITLE: MUSCULINE  
US-09-617-092A-14

Query Match 83.1% Score 774; DB 4; Length 612;

Best Local Similarity 83.2% Pred No. 4, 36-295; Matches 240; Conservative 40; Mismatch 87; Indels 10; Gaps 6;  
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MURKIN...  
OPERATING SYSTEM: PC-DOS/MS-DOS  
CURRENT APPLICATION DATA  
APPLICATION NUMBER: US/09/569-749  
CLASSIFICATION: 314  
ATtribution/Agency Information:  
REGISTRATION NUMBER: 24, 774  
REFERENCE/DOC# NUMBER: A-5264/JDN  
TELEPHONE: (415) 981-1898  
TELEFAX: (415) 366-3249  
MAILING ADDRESS:  
LEADIN: 604 amino acids  
SEQUENCE CHARACTERISTICS:  
SYNOPSIS: single  
TOPOLOGY: linear  
US-08-569-749-2  
Query Match 71.8% Score 239; DB 4; Length 604;  
Best Local Similarity 72.8% Pred No. 3, 76-219; Matches 440; Conservative 71; Mismatch 87; Indels 6; Gaps 5;  
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